Sequence Comparison A

```
RESULT 14
AAR52749
    AAR52749 standard; Protein; 355 AA.
XX
AC
    AAR52749;
XX
    30-JAN-1995 (first entry)
DT
XX
DE
    C-C chemokine receptor.
XX
KW
    C-C CKR-1; cytokine; inflammation.
XX
os
    Homo sapiens.
XX
PN
    WO9411504-A.
XX
PD
     26-MAY-1994.
XX
PF
     04-NOV-1993;
                   93WO-US10672.
XX
PR
    10-NOV-1992;
                   92US-0974025.
ХX
PA
     (GETH ) GENENTECH INC.
XX
ΡI
    Horuk R, Neote K, Schall T;
XX
DR
    WPI; 1994-183505/22.
    N-PSDB; AAQ62695.
DR
XX
PT
     New C-C chemokine receptor and nucleic acid - are used to develop
PT
    prods. for use in diagnosis and therapy of inflammation and other
PT
    cytokine-mediated disorders
XX
    Claim 1; Fig 9; 90pp; English.
PS
XX
    The sequence is that of the C-C chemokine receptor. The sequence can
CC
CC
    be used in therapeutic or diagnostic compsns. for inflammation and
CC
    other cytokine mediated disorders.
CC
    See also AAR52750-2.
XX
SQ
    Sequence
              355 AA;
 Query Match 11.5%; Score 41; DB 15; Length 355; Best Local Similarity 100.0%; Pred. No. 1.2e-31;
          41; Conservative
                                0; Mismatches
                                                  0; Indels
Qу
      115 TGLYSEIFFIILLTIDRYLAIVHAVFALRARTVTFGVITSI 155
          Db
      115 tglyseiffiilltidrylaivhavfalrartvtfgvitsi 155
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Sequence Comparison B

```
RESULT 14
AAR52749
ID
    AAR52749 standard; Protein; 355 AA.
\mathbf{x}\mathbf{x}
AC
    AAR52749;
XX
    30-JAN-1995 (first entry)
DT
XX
    C-C chemokine receptor.
DΕ
xx
KW
    C-C CKR-1; cytokine; inflammation.
XX
os
    Homo sapiens.
XX
PN
    WO9411504-A.
XX
PD
    26-MAY-1994.
XX
    04-NOV-1993;
                   93WO-US10672.
PF
XX
PR
    10-NOV-1992;
                  92US-0974025.
XX
     (GETH ) GENENTECH INC.
PA
XX
ΡI
    Horuk R, Neote K, Schall T;
XX
    WPI; 1994-183505/22.
DR
DR
    N-PSDB; AAQ62695.
ХX
PT
    New C-C chemokine receptor and nucleic acid - are used to develop
PT
    prods. for use in diagnosis and therapy of inflammation and other
PT
    cytokine-mediated disorders
XX
PS
    Claim 1; Fig 9; 90pp; English.
XX
CC
    The sequence is that of the C-C chemokine receptor. The sequence can
CC
    be used in therapeutic or diagnostic compsns. for inflammation and
CC
    other cytokine mediated disorders.
CC
    See also AAR52750-2.
XX
SQ
    Sequence
              355 AA;
                         11.5%; Score 41; DB 15; Length 355;
 Query Match
 Best Local Similarity 100.0%; Pred. No. 2.1e-32;
                               0; Mismatches
 Matches 41; Conservative
                                               0; Indels
                                                               0; Gaps
                                                                          0;
     115 TGLYSEIFFIILLTIDRYLAIVHAVFALRARTVTFGVITSI 155
         Db
     115 tglyseiffiilltidrylaivhavfalrartvtfgvitsi 155
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Sequence Comparison C

```
RESULT 13
US-08-450-393A-13
; Sequence 13, Application US/08450393A
; Patent No. 5707815
; GENERAL INFORMATION:
     APPLICANT: Charo, Israel
     APPLICANT: Coughlin, Shaun
     TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT TITLE OF INVENTION: PROTEIN RECEPTORS
     NUMBER OF SEQUENCES: 14
     CORRESPONDENCE ADDRESS:
      ADDRESSEE: Cooley Godward Castro Huddleson & Tatum
       STREET: 5 Palo Alto Square
      CITY: Palo Alto
       STATE: California
       COUNTRY: USA
       ZIP: 94306-2155
     COMPUTER READABLE FORM:
      MEDIUM TYPE: Floppy disk
       COMPUTER: IBM PC compatible
       OPERATING SYSTEM: PC-DOS/MS-DOS
       SOFTWARE: PatentIn Release #1.0, Version #1.25
     CURRENT APPLICATION DATA:
      APPLICATION NUMBER: US/08/450,393A
       FILING DATE: May 25, 1995
       CLASSIFICATION: 424
    ATTORNEY/AGENT INFORMATION:
      NAME: Cserr, Luann
      REGISTRATION NUMBER: 31,822
      REFERENCE/DOCKET NUMBER: UCAL-237/02US
     TELECOMMUNICATION INFORMATION:
      TELEPHONE: 415-843-5165
      TELEFAX: 415-8857-0663
      TELEX: 380816CooleyPA
  INFORMATION FOR SEQ ID NO: 13:
    SEQUENCE CHARACTERISTICS:
      LENGTH: 31 amino acids
      TYPE: amino acid
      TOPOLOGY: linear
    MOLECULE TYPE: peptide
US-08-450-393A-13
 Query Match 6.2%; Score 22; DB 1; Length 31; Best Local Similarity 100.0%; Pred. No. 1e-13;
 Matches 22; Conservative 0; Mismatches 0; Indels
                                                                0; Gaps
                                                                              0;
Qу
      121 IFFIILLTIDRYLAIVHAVFAL 142
          111111111111111111111
Db
        1 IFFIILLTIDRYLAIVHAVFAL 22
```